SEQUENCE LISTING

<110> Ian Popoff
Jacqueline Wyatt

<120> ANTISENSE MODULATION OF DAMAGE-SPECIFIC DNA BINDING PROTEIN 1, P127 EXPRESSION

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Val Val Thr Ala Gln Lys Pro Thr Ala Val Asn Gly Cys Val Thr Gly
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cac ttt act tcg gcc gaa gac tta aac ctg ttg att gcc aaa aac acg     211
His Phe Thr Ser Ala Glu Asp Leu Asn Leu Leu Ile Ala Lys Asn Thr
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aga tta gag atc tat gtg gtc acc gcc gag ggg ctt cgg ccc gtc aaa     259
Arg Leu Glu Ile Tyr Val Val Thr Ala Glu Gly Leu Arg Pro Val Lys
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gag gtg ggc atg tat ggg aag att gcg gtc atg gag ctt ttc agg ccc     307
Glu Val Gly Met Tyr Gly Lys Ile Ala Val Met Glu Leu Phe Arg Pro
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aag ggg gag agc aag gac ctg ctg ttt atc ttg aca gcg aag tac aat     355
Lys Gly Glu Ser Lys Asp Leu Leu Phe Ile Leu Thr Ala Lys Tyr Asn
70                  75                  80                  85

gcc tgc atc ctg gag tat aaa cag agt ggc gag agc att gac atc att     403
Ala Cys Ile Leu Glu Tyr Lys Gln Ser Gly Glu Ser Ile Asp Ile Ile
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acg cga gcc cat ggc aat gtc cag gac cgc att ggc cgc ccc tca gag     451
Thr Arg Ala His Gly Asn Val Gln Asp Arg Ile Gly Arg Pro Ser Glu
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acc ggc att att ggc atc att gac cct gag tgc cgg atg att ggc ctg     499
Thr Gly Ile Ile Gly Ile Ile Asp Pro Glu Cys Arg Met Ile Gly Leu
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cgt ctc tat gat ggc ctt ttc aag gtt att cca cta gat cgc gat aat     547
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aaa gaa ctc aag gcc ttc aac atc cgc ctg gag gag ctg cat gtc att     595
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gat gtc aag ttc cta tat ggt tgc caa gca cct act att tgc ttt gtc     643
Asp Val Lys Phe Leu Tyr Gly Cys Gln Ala Pro Thr Ile Cys Phe Val
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tac cag gac cct cag ggg cgg cac gta aaa acc tat gag gtg tct ctc     691
Tyr Gln Asp Pro Gln Gly Arg His Val Lys Thr Tyr Glu Val Ser Leu
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cga gaa aag gaa ttc aat aag ggc cct tgg aaa cag gaa aat gtc gaa     739
Arg Glu Lys Glu Phe Asn Lys Gly Pro Trp Lys Gln Glu Asn Val Glu
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ctg gct att gcc cct cct atc atc aag caa agc acg att gtg tgc cac     883
Leu Ala Ile Ala Pro Pro Ile Ile Lys Gln Ser Thr Ile Val Cys His
                250                 255                 260

aat cga gtg gac cct aat ggc tca aga tac ctg ctg gga gac atg gaa     931
Asn Arg Val Asp Pro Asn Gly Ser Arg Tyr Leu Leu Gly Asp Met Glu
            265                 270                 275

ggc cgg ctc ttc atg ctg ctt ttg gag aag gag gaa cag atg gat ggc     979
Gly Arg Leu Phe Met Leu Leu Leu Glu Lys Glu Glu Gln Met Asp Gly
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acc gtc act ctc aag gat ctc cgt gta gaa ctc ctt gga gag acc tct    1027
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att gct gag tgc ttg aca tac ctt gat aat ggt gtt gtg ttt gtc ggg    1075
Ile Ala Glu Cys Leu Thr Tyr Leu Asp Asn Gly Val Val Phe Val Gly
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tct cgc ctg ggt gac tcc cag ctt gtg aag ctc aac gtt gac agt aat    1123
Ser Arg Leu Gly Asp Ser Gln Leu Val Lys Leu Asn Val Asp Ser Asn
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gaa caa ggc tcc tat gta gtg gcc atg gaa acc ttt acc aac tta gga    1171
Glu Gln Gly Ser Tyr Val Val Ala Met Glu Thr Phe Thr Asn Leu Gly
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ccc att gtc gat atg tgc gtg gtg gac ctg gag agg cag ggg cag ggg    1219
Pro Ile Val Asp Met Cys Val Val Asp Leu Glu Arg Gln Gly Gln Gly
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cag ctg gtc act tgc tct ggg gct ttc aag gaa ggt tct ttg cgg atc    1267
Gln Leu Val Thr Cys Ser Gly Ala Phe Lys Glu Gly Ser Leu Arg Ile
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atc cgg aat gga att gga atc cac gag cat gcc agc att gac tta cca    1315
Ile Arg Asn Gly Ile Gly Ile His Glu His Ala Ser Ile Asp Leu Pro
390                 395                 400                 405

ggc atc aaa gga tta tgg cca ctg cgg tct gac cct aat cgt gag act    1363
Gly Ile Lys Gly Leu Trp Pro Leu Arg Ser Asp Pro Asn Arg Glu Thr
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gat gac act ttg gtg ctc tct ttt gtg ggc cag aca aga gtt ctc atg    1411
Asp Asp Thr Leu Val Leu Ser Phe Val Gly Gln Thr Arg Val Leu Met
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cag atc act tca gca tcg gtg agg ttg gtc tct caa gaa ccc aaa gct    1555
Gln Ile Thr Ser Ala Ser Val Arg Leu Val Ser Gln Glu Pro Lys Ala
470                 475                 480                 485

ctg gtc agt gaa tgg aag gag cct cag gcc aag aac atc agt gtg gcc    1603
Leu Val Ser Glu Trp Lys Glu Pro Gln Ala Lys Asn Ile Ser Val Ala
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tcc tgc aat agc agc cag gtg gtg gtg gct gta ggc agg gcc ctc tac    1651
Ser Cys Asn Ser Ser Gln Val Val Val Ala Val Gly Arg Ala Leu Tyr
            505                 510                 515

tat ctg cag atc cat cct cag gag ctc cgg cag atc agc cac aca gag    1699
Tyr Leu Gln Ile His Pro Gln Glu Leu Arg Gln Ile Ser His Thr Glu
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atg gaa cat gaa gtg gct tgc ttg gac atc acc cca tta gga gac agc    1747
Met Glu His Glu Val Ala Cys Leu Asp Ile Thr Pro Leu Gly Asp Ser
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aat gga ctg tcc cct ctt tgt gcc att ggc ctc tgg acg gac atc tcg    1795
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gct cgt atc ttg aag ttg ccc tct ttt gaa cta ctg cac aag gag atg    1843
Ala Arg Ile Leu Lys Leu Pro Ser Phe Glu Leu Leu His Lys Glu Met
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ctg ggt gga gag atc att cct cgc tcc atc ctg atg acc acc ttt gag    1891
Leu Gly Gly Glu Ile Ile Pro Arg Ser Ile Leu Met Thr Thr Phe Glu
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agt agc cat tac ctc ctt tgt gcc ttg gga gat gga gcg ctt ttc tac    1939
Ser Ser His Tyr Leu Leu Cys Ala Leu Gly Asp Gly Ala Leu Phe Tyr
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ttt ggg ctc aac att gag aca ggt ctg ttg agc gac cgt aag aag gtg    1987
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act ttg ggc acc cag ccc acc gta ttg agg act ttt cgt tct ctt tct    2035
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acc acc aac gtc ttt gct tgt tct gac cgc ccc act gtc atc tat agc    2083
Thr Thr Asn Val Phe Ala Cys Ser Asp Arg Pro Thr Val Ile Tyr Ser
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agc aac cac aaa ttg gtc ttc tca aat gtc aac ctc aag gaa gtg aac    2131
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gcc aac aat agc acc ctc acc att ggc acc atc gat gag atc cag aag    2227
Ala Asn Asn Ser Thr Leu Thr Ile Gly Thr Ile Asp Glu Ile Gln Lys
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ctg cac att cgc aca gtt ccc ctc tat gag tct cca agg aag atc tgc    2275
Leu His Ile Arg Thr Val Pro Leu Tyr Glu Ser Pro Arg Lys Ile Cys
710                 715                 720                 725

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Tyr Gln Glu Val Ser Gln Cys Phe Gly Val Leu Ser Ser Arg Ile Glu
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Val Gln Asp Thr Ser Gly Gly Thr Thr Ala Leu Arg Pro Ser Ala Ser
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Thr Gln Ala Leu Ser Ser Ser Val Ser Ser Ser Lys Leu Phe Ser Ser
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Ser Thr Ala Pro His Glu Thr Ser Phe Gly Glu Glu Val Glu Val His
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Gln Phe Leu Gln Asn Glu Tyr Ala Leu Ser Leu Val Ser Cys Lys Leu
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Gly Lys Asp Pro Asn Thr Tyr Phe Ile Val Gly Thr Ala Met Val Tyr
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cct gaa gag gca gag ccc aag cag ggt cgc att gtg gtc ttt cag tat    2659
Pro Glu Glu Ala Glu Pro Lys Gln Gly Arg Ile Val Val Phe Gln Tyr
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tcg gat gga aaa cta cag act gtg gct gaa aag gaa gtg aaa ggg gcc    2707
Ser Asp Gly Lys Leu Gln Thr Val Ala Glu Lys Glu Val Lys Gly Ala
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Val Tyr Ser Met Val Glu Phe Asn Gly Lys Leu Leu Ala Ser Ile Asn
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agc acg gtg cgg ctc tat gag tgg aca aca gag aag gag ctg cgc act    2803
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ggc gac ttc atc ctg gtg ggc gac ctt atg cgc tca gtg ctg ctg ctt    2899
Gly Asp Phe Ile Leu Val Gly Asp Leu Met Arg Ser Val Leu Leu Leu
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gcc tac aag ccc atg gaa gga aac ttt gaa gag att gct cga gac ttt    2947
Ala Tyr Lys Pro Met Glu Gly Asn Phe Glu Glu Ile Ala Arg Asp Phe
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